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LAW OFFICES
NIKOLAI &
MERSEREAU, P.A.

INTERNATIONAL CENTRE
900 SECOND AVENUE SOUTH, SUITE 820
MINNEAPOLIS, MINNESOTA 55402-3813
TELEPHONE 612 339-7461
FACSIMILE 612 349-6556



June 28, 2005

Our Docket No. 20020065.ORI

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

App : James Oliver Dolly et al :
 S.N. : 10/049,967 : Art Unit unknown
 Filed : February 18, 2002
 For : ISOFORMS OF SNARE MOLECULES AND THE
 USES THEREOF IN MODULATION OF
 CELLULAR EXOCYTOSIS

TRANSMITTAL LETTER

Commissioner for Patents
 P.O. Box 1450
 Alexandria, VA 22313-1450

Dear Sir:

Transmitted herewith is a Response, a substitute computer readable form (CRF), and a paper copy of the substitute Sequence Listing, in reply to the Notification of Defective Response bearing a date mailed of June 21, 2005 (copy attached), together with a copy of the marked-up Sequence Listing.

The Commissioner is authorized to charge any fees listed in 37 CFR 1.16 and 1.17 which may be required by this paper to Deposit Account No. 08-1265.

Respectfully submitted,
 NIKOLAI & MERSEREAU, P.A.


 C. G. Mersereau
 Registration No. 26205
 Attorneys for Applicant(s)
 820 International Centre
 900 Second Avenue South
 Minneapolis, MN 55402
 (612) 339-7461

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#19

PATENT APPLICATION

Our Docket No. 20020065.ORI

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Re App : James Oliver Dolly et al : June 28, 2005
S.N. : 10/049,967 : Art Unit unknown
Filed : February 18, 2002
For : ISOFORMS OF SNARE MOLECULES AND THE
USES THEREOF IN MODULATION OF
CELLULAR EXOCYTOSIS

RESPONSE

Commissioner for Patents
P.O. Box 1450
Alexandria, VA 22313-1450

Dear Sir or Madam:

This paper is submitted in response to a Notification of Defective Response dated June 21, 2005 which carried a shortened-statutory period for response of one month so that a reply is due by July 21, 2005.

Accordingly, applicants submit herewith a copy of that Notice, together with attachments, with this response.

A substitute copy of the "Sequence Listing" in computer readable form with Verification Summary Sheet is also submitted with this letter. It is believed that the content of the substitute computer readable form does comply with the requirements of 37 CFR 1.822 and/or 1.823, as all indicated

corrections have been made. A copy of the marked-up "Raw Sequence Listing" previously submitted is also attached, together with a printout of the computer readable "Sequence Listing".

The undersigned attorney hereby states that to the best of his present knowledge and belief, the information recorded in computer readable form is identical to that of the written sequence listing and includes no new matter.

The Commissioner is authorized to charge any additional fees or refund any overpayment under 37 CFR 1.16 and 1.17 which may be required by this paper to Deposit Account No. 08-1265.

Respectfully submitted,

NIKOLAI & MERSEREAU, P.A.



C. G. Mersereau
Registration No. 26205
820 International Centre
900 Second Avenue So.
Minneapolis, MN 55402
Telephone: (612) 339-7461

CERTIFICATE OF MAILING

I hereby certify that the foregoing Response, a copy of the Notification of Defective Response, a copy of the "Sequence Listing" in computer readable form (CRF) and a copy of the marked-up "Raw Sequence Listing" previously submitted, together with a printout of the computer readable "Sequence Listing", and a Verification Summary Sheet in application Serial No. 10/049,967, filed February 18, 2002, of James O. Dolly et al, entitled "ISOFORMS OF SNARE MOLECULES AND THE USES THEREOF IN MODULATION OF CELLULAR EXOCYTOSIS", are being deposited with the U.S. Postal Service as First Class mail in an envelope addressed to Mail Stop Missing Parts, Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450, postage prepaid, on June 28, 2005.



Barbara L. Davis
on behalf of C. G. Mersereau
Attorney for Applicant

Date of Signature: June 28, 2005



UNITED STATES PATENT AND TRADEMARK OFFICE

UNITED STATES DEPARTMENT OF COMMERCE
 United States Patent and Trademark Office
 Address: COMMISSIONER FOR PATENTS
 P.O. Box 1450
 Alexandria, Virginia 22313-1450
www.uspto.gov

U.S. APPLICATION NUMBER NO.	FIRST NAMED APPLICANT	ATTY. DOCKET NO.
10/049,967	James Oliver Dolly	20020065.ORI

INTERNATIONAL APPLICATION NO.

PCT/GB00/03196

I.A. FILING DATE

08/18/2000

PRIORITY DATE

C G Mersereau
 Nikolai & Mersereau
 900 Second Avenue South
 Suite 820 International Centre
 Minneapolis, MN 55402

CONFIRMATION NO. 6222
 371 FORMALITIES LETTER



OC000000016329400

Date Mailed: 06/21/2005

NOTIFICATION OF DEFECTIVE RESPONSE

The following items have been submitted by the applicant or the IB to the United States Patent and Trademark Office as a Designated / Elected Office (37 CFR 1.495)

- Copy of the International Application filed on 02/18/2002
- Copy of the International Search Report filed on 02/18/2002
- Copy of IPE Report filed on 02/18/2002
- Preliminary Amendments filed on 02/18/2002
- Information Disclosure Statements filed on 05/21/2002
- Biochemical Sequence Diskette filed on 11/04/2004
- Oath or Declaration filed on 02/23/2004
- Biochemical Sequence Listing filed on 11/04/2004
- Copy of references cited in ISR filed on 02/18/2002
- U.S. Basic National Fees filed on 02/18/2002
- Priority Documents filed on 11/04/2004

Applicant's response filed 11/04/2004 is hereby acknowledged. The following requirements set forth in the NOTIFICATION of MISSING REQUIREMENTS mailed 04/25/2002 have not been completed.

- The paper or compact disc copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 CFR 1.821(e). Applicant must provide a substitute paper or compact disc copy of the "Sequence Listing", as well as an amendment directing its entry into the application OR a substitute computer readable form (CRF) copy of the "Sequence Listing". These two items must be the same. Applicant must also provide a statement that the content of the sequence listing information recorded in computer readable form is identical to the written (on paper or compact disc) sequence listing and, where applicable, includes no new matter, as required by 37 CFR 1.821(e), 1.821(f), 1.821(g), 1.825(b), or 1.825(d). If the effective filing date is on or after September 8, 2000, see the final rulemaking notice published in the Federal Register at 65 FR 54604 (September 8, 2000) and 1238 OG 145 (September 19, 2000).
- A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 CFR 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing." Applicant must provide a substitute computer readable form (CRF) copy of the "Sequence Listing" and a statement that the content

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of the sequence listing information recorded in computer readable form is identical to the written (on paper or compact disc) sequence listing and, where applicable, includes no new matter, as required by 37 CFR 1.821(e), 1.821(f), 1.821(g), 1.825(b), or 1.825(d).

Applicant is required to complete the response within a time limit of ONE MONTH from the date of this Notification or within the time remaining in the response set forth in the Notification of Missing Requirements, whichever is the longer. No extension of this time limit may be granted under 37 CFR 1.136, but the period for response set in the Notification of Missing Requirements may be extended under 37 CFR 1.136(a).

For questions regarding compliance to 37 CFR 1.821-1.825 requirements, please contact:

- For Rules Interpretation, call (571) 272-0951
- For Patentin Software Program Help, call Patent EBC at 1-866-217-9197 or directly at 703-305-3028 / 703-308-6845 between the hours of 6 a.m. and 12 midnight, Monday through Friday, EST.
- Send e-mail correspondence for Patentin Software Program Help @ ebc@uspto.gov

Applicant is reminded that any communications to the United States Patent and Trademark Office must be mailed to the address given in the heading and include the U.S. application no. shown above (37 CFR 1.5)

A copy of this notice MUST be returned with the response.

VONDA M WALLACE

Telephone: (703) 308-9140 EXT 225

PART 1 - ATTORNEY/APPLICANT COPY

U.S. APPLICATION NUMBER NO.	INTERNATIONAL APPLICATION NO.	ATTY. DOCKET NO.
10/049,967	PCT/GB00/03196	20020065.ORI

FORM PCT/DO/EO/916 (371 Formalities Notice)

10

STIC Biotechnology Systems Branch

**RAW SEQUENCE LISTING
ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source:

Date Processed by STIC:

10/049,967C
PC
01/21/05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office; and instead should be sent via the following to the indicated addresses.

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05

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Raw Sequence Listing Error Summary

<u>ERROR DETECTED</u>	<u>SUGGESTED CORRECTION</u>	<u>SERIAL NUMBER:</u>
ATTN: NEW RULES CASES: PLEASE DISREGARD ENCLISII-ALPHA HEADERS WHICH WERE INSERTED BY PTO SOFTWARE		
1 <input checked="" type="checkbox"/> Wrapped Nucleic Wrapped Amino	The number(s) at the end of each line "wrapped" down to the next line. This may occur if your file was received in a word processor after creating it. Please adjust your right margin to .); this will prevent "wrapping."	
2 <input type="checkbox"/> Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 <input checked="" type="checkbox"/> Misaligned Amino Numbering	The numbering under each 3 ^a amino acid is misaligned. Do not use tab codes between numbers; use space characters instead.	
4 <input type="checkbox"/> Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 <input type="checkbox"/> Variable Length	Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<22> section that some may be missing.	
6 <input type="checkbox"/> PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<22> section to be missing from amino acid sequence(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<22> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<22> sections for Artificial or Unknown sequences.	
7 <input type="checkbox"/> Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence: (i) INFORMATION FOR SEQ ID NO X (insert SEQ ID NO where 'X' is shown) (ii) SEQUENCE CHARACTERISTICS (Do not insert any subheadings under this heading) (iii) SEQUENCE DESCRIPTION SEQ ID NO X (insert SEQ ID NO where 'X' is shown) This sequence is intentionally skipped. Please also adjust the '(ii) NUMBER OF SEQUENCES' response to include the skipped sequence.	
8 <input type="checkbox"/> Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence: <210> sequence id number <400> sequence id number 000	
9 <input type="checkbox"/> Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<22> is MANDATORY if n's or Xaa's are present in <220> to <22> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
10 <input type="checkbox"/> Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are Unknown, Artificial Sequence, or scientific name (GenusSpecies). <220>-<22> section is required when <213> response is Unknown or Artificial Sequence.	
11 <input type="checkbox"/> Use of <220>	Sequence(s) _____ missing the <220> feature and associated numeric identifiers and responses. Use of <220> to <22> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <22> section (See "Federal Register," 00/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
12 <input type="checkbox"/> PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 <input type="checkbox"/> Misuse of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid	

AMC - Biotechnology Systems Branch - 09/09/2003

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Page 1 of 5



PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/049,967C

DATE: 06/02/2005

TIME: 08:23:10

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\06022005\J049967C.raw

2 <110> APPLICANT: Imperial College Innovations Limited
W--> 3 <120> TITLE OF INVENTION: Isoforms of Snare Molecules and the Uses Thereof in
W--> 4 Modulation of Cellular Exocytosis
W--> 5 <130> FILE REFERENCE: ICOY/P23294PC
W--> 6 <140> CURRENT APPLICATION NUMBER: US/10/049,967C
7 <141> CURRENT FILING DATE: 2002-02-18
W--> 8 <160> NUMBER OF SEQ ID: 28
9 <170> SOFTWARE: PatentIn Ver. 2.0

ERRORED SEQUENCES

26 <210> SEQ ID NO: 3 Found 28
27 <211> LENGTH: 29
28 <212> TYPE: DNA
29 <213> ORGANISM: Artificial Sequence
W--> 30 <220> FEATURE:
31 <223> OTHER INFORMATION: Description of Artificial Sequence: pcr primer
W--> 32 <400> SEQUENCE: 3
E--> 33 ~~catcttgg ttgcgtttttat~~ 29-28

34 <210> SEQ ID NO: 4 Found 29
35 <211> LENGTH: 30
36 <212> TYPE: DNA
37 <213> ORGANISM: Artificial Sequence
W--> 38 <220> FEATURE:

39 <223> OTHER INFORMATION: Description of Artificial Sequence: pcr primer
W--> 40 <400> SEQUENCE: 4 30 delete
E--> 41 ~~catcttgg ttgcgtttttat~~ 29

42 <210> SEQ ID NO: 5 Found 31
43 <211> LENGTH: 32
44 <212> TYPE: DNA
45 <213> ORGANISM: Artificial Sequence
W--> 46 <220> FEATURE:

47 <223> OTHER INFORMATION: Description of Artificial Sequence: pcr primer
W--> 48 <400> SEQUENCE: 5
E--> 49 ~~aggatctttt ttaaccacca gtttgtttttatc~~ 32-31

162 <210> SEQ ID NO: 20
163 <211> LENGTH: 10
164 <212> TYPE: PRT
165 <213> ORGANISM: Homo Sapiens
168 <400> SEQUENCE: 20
169 Gln Arg Ala Thr Lys Met Lau Gly Ser Gly
E--> 170 1 1 5 8 10 10

Does Not Comply
Corrected Diskette Needed
(Pg. 1-2) ↗

pls group amino acids
in sets of 10s

atgcctgtgc 20
atgcctgtgc 20
aaatgtgttc 20

pls group amino
acids in sets of
10s. For ex: ↗

See item #3 on error summary
sheet.

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/049,967C

DATE: 06/02/2005
TIME: 08:23:10

Input Set : A:\PTO.DA.txt
Output Set: N:\CRF4\06022005\J049967C.raw

171 <210> SEQ ID NO: 21
 172 <211> LENGTH: 10
 173 <212> TYPE: PRT
 174 <213> ORGANISM: Artificial Sequence
 W--> 175 <220> FEATURE:
 176 <223> OTHER INFORMATION: Description of Artificial Sequence: synthesized
 W--> 177 <400> SEQUENCE: 21
 178 Gln Thr Ala Thr Lys Met Leu Gly Ser Gly
 E--> 179 | x 5 5 10 10 ✓
 216 <210> SEQ ID NO: 26
 217 <211> LENGTH: 5
 218 <212> TYPE: PRT
 219 <213> ORGANISM: Artificial Sequence
 W--> 220 <220> FEATURE:
 221 <223> OTHER INFORMATION: Description of Artificial Sequence: synthesized
 224 <400> SEQUENCE: 26
 225 Gln Arg Ala Thr Lys
 E--> 226 | x 5 5 ✓
 227 <210> SEQ ID NO: 27
 228 <211> LENGTH: 5
 229 <212> TYPE: PRT
 230 <213> ORGANISM: Artificial Sequence
 W--> 231 <220> FEATURE:
 232 <223> OTHER INFORMATION: Description of Artificial Sequence: synthesized
 W--> 233 <400> SEQUENCE: 27
 234 Gln Thr Ala Thr Lys
 E--> 235 | x 5 5 ✓

✓

See item #
on error
summary
Sheet

Best Available Copy

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/049,967C

DATE: 06/02/2005

TIME: 08:23:11

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\06022005\J049967C.raw

L:3 M:283 W: Missing Blank Line separator, <120> field identifier
L:5 M:283 W: Missing Blank Line separator, <130> field identifier
L:6 M:283 W: Missing Blank Line separator, <140> field identifier
L:8 M:283 W: Missing Blank Line separator, <160> field identifier
L:10 M:283 W: Missing Blank Line separator, <210> field identifier
L:14 M:283 W: Missing Blank Line separator, <220> field identifier
L:16 M:283 W: Missing Blank Line separator, <400> field identifier
L:22 M:283 W: Missing Blank Line separator, <220> field identifier
L:24 M:283 W: Missing Blank Line separator, <400> field identifier
L:30 M:283 W: Missing Blank Line separator, <220> field identifier
L:32 M:283 W: Missing Blank Line separator, <400> field identifier
L:33 M:254 E: No. of Bases conflict, LENGTH: Input:29 Counted:28 SEQ:3
L:33 M:252 E: No. of Seq. differs, <211> LENGTH: Input:29 Found:28 SEQ:3
L:38 M:283 W: Missing Blank Line separator, <220> field identifier
L:40 M:283 W: Missing Blank Line separator, <400> field identifier
L:41 M:254 E: No. of Bases conflict, LENGTH: Input:0 Counted:29 SEQ:4
L:41 M:252 E: No. of Seq. differs, <211> LENGTH: Input:30 Found:29 SEQ:4
L:46 M:283 W: Missing Blank Line separator, <220> field identifier
L:48 M:283 W: Missing Blank Line separator, <400> field identifier
L:49 M:254 E: No. of Bases conflict, LENGTH: Input:32 Counted:31 SEQ:5
L:49 M:252 E: No. of Seq. differs, <211> LENGTH: Input:32 Found:31 SEQ:5
L:54 M:283 W: Missing Blank Line separator, <220> field identifier
L:56 M:283 W: Missing Blank Line separator, <400> field identifier
L:62 M:283 W: Missing Blank Line separator, <220> field identifier
L:64 M:283 W: Missing Blank Line separator, <400> field identifier
L:70 M:283 W: Missing Blank Line separator, <220> field identifier
L:72 M:283 W: Missing Blank Line separator, <400> field identifier
L:78 M:283 W: Missing Blank Line separator, <220> field identifier
L:80 M:283 W: Missing Blank Line separator, <400> field identifier
L:86 M:283 W: Missing Blank Line separator, <220> field identifier
L:88 M:283 W: Missing Blank Line separator, <400> field identifier
L:94 M:283 W: Missing Blank Line separator, <220> field identifier
L:96 M:283 W: Missing Blank Line separator, <400> field identifier
L:102 M:283 W: Missing Blank Line separator, <220> field identifier
L:104 M:283 W: Missing Blank Line separator, <400> field identifier
L:110 M:283 W: Missing Blank Line separator, <220> field identifier
L:112 M:283 W: Missing Blank Line separator, <400> field identifier
L:118 M:283 W: Missing Blank Line separator, <220> field identifier
L:120 M:283 W: Missing Blank Line separator, <400> field identifier
L:126 M:283 W: Missing Blank Line separator, <220> field identifier
L:128 M:283 W: Missing Blank Line separator, <400> field identifier
L:134 M:283 W: Missing Blank Line separator, <220> field identifier
L:136 M:283 W: Missing Blank Line separator, <400> field identifier
L:142 M:283 W: Missing Blank Line separator, <220> field identifier
L:144 M:283 W: Missing Blank Line separator, <400> field identifier
L:150 M:283 W: Missing Blank Line separator, <220> field identifier
L:152 M:283 W: Missing Blank Line separator, <400> field identifier
L:158 M:283 W: Missing Blank Line separator, <220> field identifier

Best Available Copy

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/049,967C

DATE: 06/02/2005
TIME: 08:23:11

Input Set : A:\PTO.DA.txt
Output Set: N:\CRF4\06022005\J049967C.raw

L:160 M:283 W: Missing Blank Line separator, <400> field identifier
L:170 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:20 ✓
L:175 M:283 W: Missing Blank Line separator, <220> field identifier
L:177 M:283 W: Missing Blank Line separator, <400> field identifier
L:179 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:21 ✓
L:184 M:283 W: Missing Blank Line separator, <220> field identifier
L:186 M:283 W: Missing Blank Line separator, <400> field identifier
L:193 M:283 W: Missing Blank Line separator, <220> field identifier
L:195 M:283 W: Missing Blank Line separator, <400> field identifier
L:202 M:283 W: Missing Blank Line separator, <220> field identifier
L:226 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:26 ✓
L:235 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:27 ✓

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PAGE:
06/27/2005
14:55:49

VERIFICATION SUMMARY REPORT
PATENT APPLICATION

DATE:
TIME:

INPUT SEQ: C:\Documents and Settings\barbara
\Desktop\20020065-2Seq.List.txt

GENERAL INFORMATION SECTION

3,<110> Imperial College Innovations Limited
5,<120> Isoforms of Snare Molecules and the Uses Thereof in
7, Modulation of Cellular Exocytosis
9,<130> ICOY/P23294PC
11,<140> US/10/049,967
12,<141> 2002-02-18
14,<160> 28
15,<170> PatentIn Ver. 2.0

ERRORED LINES SECTION

STATISTICS SUMMARY

Application Serial Number: US/10/049,967
Alpha or Numeric or Xml: Numeric
Application Class:
Application File Date: 2002-02-18
Art Unit:
Software Application: PatentIN2.0
Total Number of Sequences: 28
Total Nucleotides: 665
Total Amino Acids: 60
Number of Errors: 0
Number of Warnings: 0
Number of Corrections: 0